TECH CENTER 1600/2900

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<110> Estell, David A.
Harding, Fiona A.
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<120> PROTEINS PRODUCING AN ALTERED IMMUNOGENIC RESPONSE AND METHODS OF MAKING AND USING THE SAME

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<140> US 09/500,135
<141> 2000-02-08
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<150> US 09/060,872 <151> 1998-04-15

<160> 236

<170> PatentIn Ver. 2.1

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<210> 1
<211> 1495
<212> DNA
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<213> Bacillus amyloliquefaciens

<220>

<221> mat\_peptide <222> (417)..(1495)

<220>

<221> CDS

<222> (96)..(1244)

<220>

<221> misc\_feature

<222> (582)..(584)

<223> The nnn at positions 582 through 584 which in a preferred embodiment (aat) is to code for asparagine, but which may also code for proline.

<220>

<221> misc\_feature

<222> (585)..(587)

<223> The nnn at positions 585 through 587 which in a
 preferred embodiment (cct) is to code for proline,
 but which may also code for asparagine.

<220>

<221> misc\_feature

<222> (597)..(599)

<223> The nnn at positions 597 to 599 which in a
 preferred embodiment (aac) is to code for
 asparagine, but which may also code for aspartic acid.

<220>

<221> misc feature

<222> (678)..(680)

<223> The nnn at positions 678 through 680 which in a preferred embodiment (gca) is to code for alanine, but which may also code for serine. <220> <221> misc feature <222> (681)..(683) <223> The nnn at positions 681 through 683 which in a preferred embodiment (tca) is to code for serine, but which may also code for alanine. <220> <221> misc\_feature <222> (708)..(710) <223> The nnn at positions 708 through 710 which in a preferred embodiment (gct) is to code for alanine, but which may also code for aspartic acid. <220> <221> misc feature <222> (711)..(713) <223> The nnn at positions 711 through 713 which in a preferred embodiment (gac) is to code for aspartic acid, but which may also code for alanine. <220> <221> misc\_feature <222> (888)..(890) <223> The nnn at positions 888 through 890 which in a preferred embodiment (act) is to code for threonine, but which may also code for serine. <220> <221> misc feature <222> (891)..(893) <223> The nnn at positions 891 through 893 which in a preferred embodiment (tcc) is to code for serine, but which may also code for threonine. <220> <221> misc feature <222> (1167)..(1169) <223> The nnn at positions 1167 through 1169 which in a preferred embodiment (gaa) is to code for glutamic acid, but which may also code for glutamine. ggtctactaa aatattattc catactatac aattaataca cagaataatc tgtctattgg 60

ttattctgca aatgaaaaaa aggagaggat aaaga atg aga ggc aaa aaa gta 113 Met Arg Gly Lys Lys Val -105

tgg atc agt ttg ctg ttt gct tta gcg tta atc ttt acg atg gcg ttc 161
Trp Ile Ser Leu Leu Phe Ala Leu Ala Leu Ile Phe Thr Met Ala Phe
-100 -95 -90

				tct Ser												209
aaa Lys	tat Tyr	att Ile	gtc Val	999 Gly -65	ttt Phe	aaa Lys	cag Gln	aca Thr	atg Met -60	agc Ser	acg Thr	atg Met	agc Ser	gcc Ala -55	gct Ala	257
				gtc Val												305
ttc Phe	aaa Lys	tat Tyr -35	gta Val	gac Asp	gca Ala	gct Ala	tca Ser -30	gct Ala	aca Thr	tta Leu	aac Asn	gaa Glu -25	aaa Lys	gct Ala	gta Val	353
aaa Lys	gaa Glu -20	ttg Leu	aaa Lys	aaa Lys	gac Asp	ccg Pro -15	agc Ser	gtc Val	gct Ala	tac Tyr	gtt Val -10	gaa Glu	gaa Glu	gat Asp	cac His	401
gta Val -5	gca Ala	cat His	gcg Ala	tac Tyr -1	gcg Ala 1	cag Gln	tcc Ser	gtg Val	cct Pro 5	tac Tyr	ggc Gly	gta Val	tca Ser	caa Gln 10	att Ile	449
				ctg Leu												497
				gac Asp												545
gta Val	gca Ala 45	ggc Gly	gga Gly	gcc Ala	agc Ser	atg Met 50	gtt Val	cct Pro	tct Ser	gaa Glu	aca Thr 55	nnn Xaa	nnn Xaa	ttc Phe	caa Gln	593
				cac His												641
aat Asn	aac Asn	tca Ser	atc Ile	ggt Gly 80	gta Val	tta Leu	ggc Gly	gtt Val	gcg Ala 85	cca Pro	agc Ser	nnn Xaa	nnn Xaa	ctt Leu 90	tac Tyr	689
				ctc Leu												737
att Ile	aac Asn	gga Gly 110	atc Ile	gag Glu	tgg Trp	gcg Ala	atc Ile 115	gca Ala	aac Asn	aat Asn	atg Met	gac Asp 120	gtt Val	att Ile	aac Asn	785
				gga Gly												833
gat	aaa	gcc	gtt	gca	tcc	ggc	gtc	gta	gtc	gtt	gcg	gca	gcc	ggt	aac	881

Asp Lys Ala Val Ala Ser Gly Val Val Val Ala Ala Ala Gly Asn 140 145 150 155													
gaa ggc nnn nnn ggc agc tca agc aca gtg ggc tac cct ggt aaa tac 929 Glu Gly Xaa Xaa Gly Ser Ser Ser Thr Val Gly Tyr Pro Gly Lys Tyr 160 165 170													
cct tct gtc att gca gta ggc gct gtt gac agc agc aac caa aga gca 977 Pro Ser Val Ile Ala Val Gly Ala Val Asp Ser Ser Asn Gln Arg Ala 175 180 185													
tet tte tea age gta gga eet gag ett gat gte atg gea eet gge gta 1029 Ser Phe Ser Ser Val Gly Pro Glu Leu Asp Val Met Ala Pro Gly Val 190 195 200	ō												
tct atc caa agc acg ctt cct gga aac aaa tac ggg gcg tac aac ggt 1073 Ser Ile Gln Ser Thr Leu Pro Gly Asn Lys Tyr Gly Ala Tyr Asn Gly 205 210 215	3												
acg tca atg gca tct ccg cac gtt gcc gga gcg gct gct ttg att ctt  Thr Ser Met Ala Ser Pro His Val Ala Gly Ala Ala Ala Leu Ile Leu 220 225 230 235	1												
tct aag cac ccg aac tgg aca aac act caa gtc cgc agc agt tta nnn 1169 Ser Lys His Pro Asn Trp Thr Asn Thr Gln Val Arg Ser Ser Leu Xaa 240 245 250	Э												
aac acc act aca aaa ctt ggt gat tct ttc tac tat gga aaa ggg ctg 121' Asn Thr Thr Lys Leu Gly Asp Ser Phe Tyr Tyr Gly Lys Gly Leu 255 260 265	7												
atc aac gta cag gcg gca gct cag taa aacataaaaa accggccttg Ile Asn Val Gln Ala Ala Gln 270 275													
gccccgccgg tttttttatt tttcttcctc cgcatgttca atccgctcca taatcgacgg 1													
atggctccct ctgaaaattt taacgagaaa cggcgggttg acccggctca gtcccgtaac 138													
ggccaagtcc tgaaacgtct caatcgccgc ttcccggttt ccggtcagct caatgccgta 1													
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Ile Phe Thr Met Ala Phe Gly Ser Thr Ser Ser Ala Gln Ala Ala Gly
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Lys Ser Asn Gly Glu Lys Lys Tyr Ile Val Gly Phe Lys Gln Thr Met
                              40
Ser Thr Met Ser Ala Ala Lys Lys Lys Asp Val Ile Ser Glu Lys Gly
                          55
Gly Lys Val Gln Lys Gln Phe Lys Tyr Val Asp Ala Ala Ser Ala Thr
                                          75
                     70
Leu Asn Glu Lys Ala Val Lys Glu Leu Lys Lys Asp Pro Ser Val Ala
                                      90
Tyr Val Glu Glu Asp His Val Ala His Ala Tyr Ala Gln Ser Val Pro
                                 105
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Tyr Gly Val Ser Gln Ile Lys Ala Pro Ala Leu His Ser Gln Gly Tyr 120 Thr Gly Ser Asn Val Lys Val Ala Val Ile Asp Ser Gly Ile Asp Ser 135 Ser His Pro Asp Leu Lys Val Ala Gly Gly Ala Ser Met Val Pro Ser 155 150 Glu Thr Xaa Xaa Phe Gln Asp Xaa Asn Ser His Gly Thr His Val Ala 165 170 Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val Ala 185 180 Pro Ser Xaa Xaa Leu Tyr Ala Val Lys Val Leu Gly Xaa Xaa Gly Ser 200 Gly Gln Tyr Ser Trp Ile Ile Asn Gly Ile Glu Trp Ala Ile Ala Asn 215 220 Asn Met Asp Val Ile Asn Met Ser Leu Gly Gly Pro Ser Gly Ser Ala 235 230 Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala Ser Gly Val Val Val 250 255 245 Val Ala Ala Gly Asn Glu Gly Xaa Xaa Gly Ser Ser Ser Thr Val 270 265 Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala Val Gly Ala Val Asp 280 Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val Gly Pro Glu Leu Asp 300 295 Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr Leu Pro Gly Asn Lys 315 310 Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser Pro His Val Ala Gly 330 325 Ala Ala Leu Ile Leu Ser Lys His Pro Asn Trp Thr Asn Thr Gln 340 345 350 Val Arg Ser Ser Leu Xaa Asn Thr Thr Thr Lys Leu Gly Asp Ser Phe 355 360 Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala Ala Gln 375

<210> 3 <211> 275

<212> PRT

<213> Bacillus amyloliquefaciens

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Ala Gln Ser Val Pro Tyr Gly Val Ser Gln Ile Lys Ala Pro Ala Leu

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His Ser Gln Gly Tyr Thr Gly Ser Asn Val Lys Val Ala Val Ile Asp 20 25 30

Ser Gly Ile Asp Ser Ser His Pro Asp Leu Lys Val Ala Gly Gly Ala 35 40 45

Ser Met Val Pro Ser Glu Thr Asn Pro Phe Gln Asp Asn Asn Ser His
50 55 60

Gly Thr His Val Ala Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly 65 70 75 80 Val Leu Gly Val Ala Pro Ser Ala Ser Leu Tyr Ala Val Lys Val Leu 85 90 95

Gly Ala Asp Gly Ser Gly Gln Tyr Ser Trp Ile Ile Asn Gly Ile Glu 100 105 110

Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn Met Ser Leu Gly Gly 115 120 125

Pro Ser Gly Ser Ala Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala 130 135 140

Ser Gly Val Val Val Val Ala Ala Ala Gly Asn Glu Gly Thr Ser Gly 145 150 155 160

Ser Ser Ser Thr Val Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala 165 170 175

Val Gly Ala Val Asp Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val 180 185 190

Gly Pro Glu Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr 195 200 205

Leu Pro Gly Asn Lys Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser 210 215 220

Pro His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Asn 225 230 235 235

Trp Thr Asn Thr Gln Val Arg Ser Ser Leu Glu Asn Thr Thr Thr Lys 245 250 255

Leu Gly Asp Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala 260 265 270

Ala Ala Gln 275

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<211> 275

<212> PRT

<213> Bacillus subtilis

<400> 4

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His Ser Gln Gly Tyr Thr Gly Ser Asn Val Lys Val Ala Val Ile Asp 20 25 30

Ser Gly Ile Asp Ser Ser His Pro Asp Leu Asn Val Arg Gly Gly Ala

Ser Phe Val Pro Ser Glu Thr Asn Pro Tyr Gln Asp Gly Ser Ser His

50					55					60				
Gly Thr 65	His	Val	Ala	Gly 70	Thr	Ile	Ala	Ala	Leu 75	Asn	Asn	Ser	Ile	Gly 80
Val Leu	Gly	Val	Ser 85	Pro	Ser	Ala	Ser	Leu 90	Tyr	Ala	Val	Lys	Val 95	Leu
Asp Ser	Thr	Gly 100	Ser	Gly	Gln	Tyr	Ser 105	Trp	Ile	Ile	Asn	Gly 110	Ile	Glu
Trp Ala	Ile 115	Ser	Asn	Asn	Met	Asp 120	Val	Ile	Asn	Met	Ser 125	Leu	Gly	Gly
Pro Thr 130	Gly	Ser	Thr	Ala	Leu 135	Lys	Thr	Val	Val	Asp 140	Lys	Ala	Val	Ser
Ser Gly 145	Ile	Val	Val	Ala 150	Ala	Ala	Ala	Gly	Asn 155	Glu	Gly	Ser	Ser	Gly 160
Ser Thr	Ser	Thr	Val 165	Gly	Tyr	Pro	Ala	Lys 170	Tyr	Pro	Ser	Thr	Ile 175	Ala
Val Gly	Ala	Val 180	Asn	Ser	Ser	Asn	Gln 185	Arg	Ala	Ser	Phe	Ser 190	Ser	Ala
Gly Ser	Glu 195	Leu	Asp	Val	Met	Ala 200	Pro	Gly	Val	Ser	Ile 205	Gln	Ser	Thr
Leu Pro 210	Gly	Gly	Thr	Tyr	Gly 215	Ala	Tyr	Asn	Gly	Thr 220	Ser	Met	Ala	Thr
Pro His 225	Val	Ala	Gly	Ala 230	Ala	Ala	Leu	Ile	Leu 235	Ser	Lys	His	Pro	Thr 240
Trp Thr	Asn	Ala	Gln 245	Val	Arg	Asp	Arg	Leu 250	Glu	Ser	Thr	Ala	Thr 255	Tyr
Leu Gly	Asn	Ser 260	Phe	Tyr	Tyr	Gly	Lys 265	Gly	Leu	Ile	Asn	Val 270	Gln	Ala
Ala Ala	Gln 275													
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Gln Ala Gln Gly Phe Lys Gly Ala Asn Val Lys Val Ala Val Leu Asp 20 25 30 Thr Gly Ile Gln Ala Ser His Pro Asp Leu Asn Val Val Gly Gly Ala 35 40 45

Ser Phe Val Ala Gly Glu Ala Tyr Asn Thr Asp Gly Asn Gly His Gly 50 55 60

Thr His Val Ala Gl; Thr Val Ala Ala Leu Asp Asn Thr Thr Gly Val 65 70 75 80

Leu Gly Val Ala Pro Ser Val Ser Leu Tyr Ala Val Lys Val Leu Asn 85 90 95

Ser Ser Gly Ser Gl; Ser Tyr Ser Gly Ile Val Ser Gly Ile Glu Trp 100 105 110

Ala Thr Thr Asn Gly Met Asp Val Ile Asn Met Ser Leu Gly Gly Ala 115 120 125

Ser Gly Ser Thr Ala Met Lys Gln Ala Val Asp Asn Ala Tyr Ala Arg 130 135 140

Gly Val Val Val Ala Ala Ala Gly Asn Ser Gly Asn Ser Gly Ser 145 150 155 160

Thr Asn Thr Ile Gly Tyr Pro Ala Lys Tyr Asp Ser Val Ile Ala Val 165 170 175

Gly Ala Val Asp Ser Asn Ser Asn Arg Ala Ser Phe Ser Ser Val Gly
180 185 190

Ala Glu Leu Glu Val Met Ala Pro Gly Ala Gly Val Tyr Ser Thr Tyr 195 200 205

Pro Thr Asn Thr Tyr Ala Thr Leu Asn Gly Thr Ser Met Ala Ser Pro 210 215 220

His Val Ala Gly Ala Ala Ala Leu Ile Leu Ser Lys His Pro Asn Leu 225 230 235 240

Ser Ala Ser Gln Val Arg Asn Arg Leu Ser Ser Thr Ala Thr Tyr Leu 245 250 255

Gly Ser Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Glu Ala Ala 260 265 270

Ala Gln

<210> 6

<211> 269

<212> PRT

<213> Human

<400> 6

Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala

1 5 10 15

His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp 20 25 30

Thr Gly Ile Ser Thr His Pro Asp Leu Asn Ile Arg Gly Gly Ala Ser 35 40 45

Phe Val Pro Gly Glu Pro Ser Thr Gln Asp Gly Asn Gly His Gly Thr 50 55 60

His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly Val Leu 65 70 75 80

Gly Val Ala Pro Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala 85 90 95

Ser Gly Ser Gly Ser Val Ser Ser Ile Ala Gl<br/>n Gly Leu Glu Trp Ala 100 105 110

Gly Asn Asn Gly Met His Val Ala Asn Leu Ser Leu Gly Ser Pro Ser 115 120 125

Pro Ser Ala Thr Leu Glu Gln Ala Val Asn Ser Ala Thr Ser Arg Gly 130 135 140

Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly Ala Thr Asp Gln
165 170 175

Asn Asn Asn Arg Ala Ser Phe Ser Gln Tyr Gly Ala Gly Leu Asp Ile 180 185 190

Val Ala Pro Gly Val Asn Val Gln Ser Thr Tyr Pro Gly Ser Thr Tyr 195 200 205

Ala Ser Leu Asn Gly Thr Ser Met Ala Thr Pro His Val Ala Gly Ala 210 215 220

Ala Ala Leu Val Lys Gln Lys Asn Pro Ser Trp Ser Asn Val Gln Ile 225 230 235 240

Arg Asn His Leu Lys Asn Thr Ala Thr Ser Leu Gly Ser Thr Asn Leu 245 250 255

Tyr Gly Ser Gly Leu Val Asn Ala Glu Ala Ala Thr Arg 260 265

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<212> PRT

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<220>

<223> Description of Artificial Sequence: Synthetic

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Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala
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Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala His Asn
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Gly Ile Ser Arg Val Gln Ala Pro Ala Ala His Asn Arg Gly Leu
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<210> 13
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Ala Pro Ala Ala His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys
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<210> 14
<211> 15
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Ala His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val
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Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp Thr
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Val Ala Val Leu Asp Thr Gly Ile Ser Thr His Pro Asp Leu Asn
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<210> 19
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Gly Ile Ser Thr His Pro Asp Leu Asn Ile Arg Gly Gly Ala Ser
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Leu Asn Asn Ser Ile Gly Val Leu Gly Val Ala Pro Ser Ala Glu
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Val Lys Val Leu Gly Ala Ser Gly Ser Gly Ser Val Ser Ser Ile
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Leu Glu Trp Ala Gly Asn Asn Gly Met His Val Ala Asn Leu Ser
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<210> 46
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Ala Gly Asn Asn Gly Met His Val Ala Asn Leu Ser Leu Gly Ser
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Asn Gly Met His Val Ala Asn Leu Ser Leu Gly Ser Pro Ser Pro
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Asn Leu Ser Leu Gly Ser Pro Ser Pro Ser Ala Thr Leu Glu Gln
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<400> 50
Leu Gly Ser Pro Ser Pro Ser Ala Thr Leu Glu Gln Ala Val Asn
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Pro Ser Pro Ser Ala Thr Leu Glu Gln Ala Val Asn Ser Ala Thr
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Ser Ala Thr Leu Glu Gln Ala Val Asn Ser Ala Thr Ser Arq Gly
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Leu Glu Gln Ala Val Asn Ser Ala Thr Ser Arg Gly Val Leu Val
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<400> 54
Ala Val Asn Ser Ala Thr Ser Arg Gly Val Leu Val Val Ala Ala
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<210> 55
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<212> PRT
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Ser Ala Thr Ser Arg Gly Val Leu Val Val Ala Ala Ser Gly Asn
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Ser Gly Ala Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala
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<210> 61
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Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val
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<210> 62
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Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly Ala Thr
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<223> Description of Artificial Sequence: Synthetic

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Asn Asn Arg Ala Ser Phe Ser Gln Tyr Gly Ala Gly Leu Asp Ile 1 10

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Ala Ser Phe Ser Gln Tyr Gly Ala Gly Leu Asp Ile Val Ala Pro
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<400> 70
Ser Gln Tyr Gly Ala Gly Leu Asp Ile Val Ala Pro Gly Val Asn
1 5
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<400> 71
Gly Ala Gly Leu Asp Ile Val Ala Pro Gly Val Asn Val Gln Ser
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<210> 72
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<400> 72
Leu Asp Ile Val Ala Pro Gly Val Asn Val Gln Ser Thr Tyr Pro
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Val Ala Pro Gly Val Asn Val Gln Ser Thr Tyr Pro Gly Ser Thr
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Gly Val Asn Val Gln Ser Thr Tyr Pro Gly Ser Thr Tyr Ala Ser
                                     10
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<210> 75
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Val Gln Ser Thr Tyr Pro Gly Ser Thr Tyr Ala Ser Leu Asn Gly
                 5
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Thr Tyr Pro Gly Ser Thr Tyr Ala Ser Leu Asn Gly Thr Ser Met
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Gly Ser Thr Tyr Ala Ser Leu Asn Gly Thr Ser Met Ala Thr Pro 1 5 10 <210> 78 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic <400> 78 Tyr Ala Ser Leu Asn Gly Thr Ser Met Ala Thr Pro His Val Ala 1 5 <210> 79 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic Leu Asn Gly Thr Ser Met Ala Thr Pro His Val Ala Gly Ala Ala 1 5 10 15 <210> 80 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic <400> 80 Thr Ser Met Ala Thr Pro His Val Ala Gly Ala Ala Ala Leu Val 5 10 <210> 81 <211> 15 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Synthetic <400> 81 Ala Thr Pro His Val Ala Gly Ala Ala Ala Leu Val Lys Gln Lys

<210> 82

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Gly Val Ala Gly Ala Ala Ala Leu Val Lys Gln Lys Asn Pro Ser 1 5

<210> 83

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<223> Description of Artificial Sequence: Synthetic

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Gly Ala Ala Leu Val Lys Gln Lys Asn Pro Ser Trp Ser Asn 5

<210> 84

<211> 15

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<400> 84

Ala Leu Val Lys Gln Lys Asn Pro Ser Trp Ser Asn Val Gln Ile 5 10

<210> 85

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<213> Artificial Sequence

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<223> Description of Artificial Sequence: Synthetic

<400> 85

Lys Gln Lys Asn Pro Ser Trp Ser Val Asn Gln Ile Arg Asn His 5 10

<210> 86

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Trp Ser Asn Val Gln Ile Arg Asn His Leu Lys Asn Thr Ala Thr
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Val Gln Ile Arg Asn His Leu Lys Asn Thr Ala Thr Ser Leu Gly
                5
                                    10
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<400> 89
Arg Asn His Leu Lys Asn Thr Ala Thr Ser Leu Gly Ser Thr Asn
1 5
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Leu Lys Asn Thr Ala Thr Ser Leu Gly Ser Thr Asn Leu Tyr Gly
                  5
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<210> 91
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Thr Ala Thr Ser Leu Gly Ser Thr Asn Leu Tyr Gly Ser Gly Leu
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Ser Leu Gly Ser Thr Asn Leu Tyr Gly Ser Gly Leu Val Asn Ala
                                    10
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<211> 15
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Ser Thr Asn Leu Tyr Gly Ser Gly Leu Val Asn Ala Glu Ala Ala
                 5
                                                         15
1
                                     10
<210> 94
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Asn Leu Tyr Gly Ser Gly Leu Val Asn Ala Glu Ala Ala Thr Arg
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Asp Ala Glu Leu His Ile Phe Arg Val Phe Thr Asn Asn Gln Val
                                    10
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Pro Leu Arg Arg Ala Ser Leu Ser Leu Gly Ser Gly Phe Trp His
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Arg Ala Ser Leu Ser Leu Gly Ser Gly Phe Trp His Ala Thr Gly
                 5
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Leu Ser Leu Gly Ser Gly Phe Trp His Ala Thr Gly Arg His Ser
                  5
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Gly Ser Gly Phe Trp His Ala Thr Gly Arg His Ser Ser Arg Arg
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Phe Trp His Ala Thr Gly Arg His Ser Ser Arg Arg Leu Leu Arg
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Ala Thr Gly Arg His Ser Ser Arg Arg Leu Leu Arg Ala Ile Pro
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Arg His Ser Ser Arg Arg Leu Leu Arg Ala Ile Pro Arg Gln Val
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                                    10
<210> 103
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Ser Arg Arg Leu Leu Arg Ala Ile Pro Arg Gln Val Ala Gln Thr
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Leu Leu Arg Ala Ile Pro Arg Gln Val Ala Gln Thr Leu Gln Ala
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Ala Ile Pro Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu
                                    10
                 5
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<211> 15
<212> PRT
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<400> 106
Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu Trp Gln Met
             5
                                    10
<210> 107
<211> 15
<212> PRT
<213> Artificial Sequence
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Ala Gln Thr Leu Gln Ala Asp Val Leu Trp Gln Met Gly Tyr Thr
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<210> 108
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<400> 108
Leu Gln Ala Asp Val Leu Trp Gln Met Gly Tyr Thr Gly Ala Asn
                                    10
<210> 109
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Asp Val Leu Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val
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Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val Ala Val Phe
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Gly Tyr Thr Gly Ala Asn Val Arg Val Ala Val Phe Asp Thr Gly
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<212> PRT
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Gly Ala Asn Val Arg Val Ala Val Phe Asp Thr Gly Leu Ser Glu
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Lvs His Pro His Phe Lys Asn Val Lys Glu Arg Thr Asn Trp Thr
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His Phe Lys Asn Val Lys Glu Arg Thr Asn Trp Thr Asn Glu Arg
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Asn Val Lys Glu Arg Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp
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Glu Arg Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp Asp Gly Leu
                                     10
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<223> Description of Artificial Sequence: Synthetic
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Asn Trp Thr Asn Glu Arg Thr Leu Asp Asp Gly Leu Gly His Gly
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<212> PRT
<213> Artificial Sequence
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<400> 122
Asn Glu Arg Thr Leu Asp Asp Gly Leu Gly His Gly Thr Phe Val
                5
<210> 123
<211> 15
<212> PRT
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Thr Leu Asp Asp Gly Leu Gly His Gly Thr Phe Val Ala Gly Val
                5
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<210> 124
<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
<400> 124
Asp Gly Leu Gly His Gly Thr Phe Val Ala Gly Val Ile Ala Ser
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<212> PRT
<213> Artificial Sequence
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Gly His Gly Thr Phe Val Ala Gly Val Ile Ala Ser Met Arg Glu
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Thr Phe Val Ala Gly Val Ile Ala Ser Met Arg Glu Cys Gln Gly
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<210> 127
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<212> PRT
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<400> 127
Ala Gly Val Ile Ala Ser Met Arg Glu Cys Gln Gly Phe Ala Pro
1 5
                      10
<210> 128
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<212> PP.T
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Ile Ala Ser Met Arg Glu Cys Gln Gly Phe Ala Pro Asp Ala Glu
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Met Arg Glu Cys Gln Gly Phe Ala Pro Asp Ala Glu Leu His Ile
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<210> 130
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Cys Gln Gly Phe Ala Pro Asp Ala Glu Leu His Ile Phe Arg Val
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Phe Ala Pro Asp Ala Glu Leu His Ile Phe Arg Val Phe Thr Asn
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<212> PRT
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Asp Ala Glu Leu His Ile Phe Arg Val Phe Thr Asn Asn Gln Val
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Leu His Ile Phe Arg Val Phe Thr Asn Asn Gln Val Ser Tyr Thr
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Phe Asn Tyr Ala Ile Leu Lys Lys Ile Asp Val Leu Asn Leu Ser
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<210> 141
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Ala Ile Leu Lys Lys Ile Asp Val Leu Asn Leu Ser Ile Gly Gly
               5
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Lys Lys Ile Asp Val Leu Asn Leu Ser Ile Gly Gly Pro Asp Phe
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Asp Val Leu Asn Leu Ser Ile Gly Gly Pro Asp Phe Met Asp His
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                                    10
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Asn Leu Ser Ile Gly Gly Pro Asp Phe Met Asp His Pro Phe Val
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Ile Gly Gly Pro Asp Phe Met Asp His Pro Phe Val Asp Lys Val
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Pro Asp Phe Met Asp His Pro Phe Val Asp Lys Val Trp Glu Leu
1 5
                                   10
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Met Asp His Pro Phe Val Asp Lys Val Trp Glu Leu Thr Ala Asn
                 5
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<400> 148
Pro Phe Val Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val Ile
              5
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<400> 149
Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val Ile Met Val Ser
                5
                                   10
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Trp Glu Leu Thr Ala Asn Asn Val Ile Met Val Ser Ala Ile Gly
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                                                        15
                5
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Thr Ala Asn Asn Val Ile Met Val Ser Ala Ile Gly Asn Asp Gly
          5
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Asn Val Ile Met Val Ser Ala Ile Gly Asn Asp Gly Pro Leu Tyr
                                    10
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Met Val Ser Ala Ile Gly Asn Asp Gly Pro Leu Tyr Gly Thr Ile
                                    10
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                 5
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<400> 154
Ala Ile Gly Asn Asp Gly Pro Leu Tyr Gly Thr Leu Asn Asn Pro
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<210> 156
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Pro Leu Tyr Gly Thr Leu Asn Asn Pro Ala Asp Gln Met Asp Val
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Gly Thr Leu Asn Asn Pro Ala Asp Gln Met Asp Val Ile Gly Val
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Asn Asn Pro Ala Asp Gln Met Asp Val Ile Gly Val Gly Gly Ile
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Ala Asp Gln Met Asp Val Ile Gly Val Gly Gly Ile Asp Phe Glu
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Asp Phe Glu Asp Asn Ile Ala Arg Phe Ser Ser Arg Gly Met Thr
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Asp Asn Ile Ala Arg Phe Ser Ser Arg Gly Met Thr Thr Trp Glu

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<210> 165

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Gly Met Thr Trp Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys
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Thr Trp Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys Pro Asp Ile
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Leu Pro Gly Gly Tyr Gly Arg Met Lys Pro Asp Ile Val Thr Tyr
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Gly Ala Gly Val Arg Gly Ser Gly Val Lys Gly Gly Cys Arg Ala
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Val Arg Gly Ser Gly Val Lys Gly Gly Cys Arg Ala Leu Ser Gly
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Ser Gly Val Lys Gly Gly Cys Arg Ala Leu Ser Gly Thr Ser Val
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Lys Gly Gly Cys Arg Ala Leu Ser Gly Thr Ser Val Ala Ser Pro
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Cys Arg Ala Leu Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala
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Leu Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala Gly Ala Val
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Val Ser Thr Val Gln Lys Arg Glu Leu Val Asn Pro Ala Ser Met
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Val Gln Lys Arq Glu Leu Val Asn Pro Ala Ser Met Lys Gln Ala
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Arg Glu Leu Val Asn Pro Ala Ser Met Lys Gln Ala Leu Ile Ala
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Val Asn Pro Ala Ser Met Lys Gln Ala Leu Ile Ala Ser Ala Arg
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Ala Ser Met Lys Gln Ala Leu Ile Ala Ser Ala Arg Arg Leu Pro
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<211> 15

<212> PRT

<213> Artificial Sequence

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<220>

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<211> 15
<212> PRT
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<400> 201
Asn Ser Tyr Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp
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<211> 15
<212> PRT
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<223> Description of Artificial Sequence: Synthetic
<400> 202
Lys Pro Gln Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu
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<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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Ala Ser Leu Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr
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<210> 204
<211> 15
<212> PRT
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Ser Pro Ser Tyr Ile Asp Leu Thr Glu Cys Pro Tyr Met Trp Pro
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<211> 15
<212> PRT
<213> Artificial Sequence
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<223> Description of Artificial Sequence: Synthetic
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Tyr Ile Asp Leu Thr Glu Cys Pro Tyr Met Trp Pro Tyr Cys Ser
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<210> 206
<211> 15
<212> PRT
<213> Artificial Sequence
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<400> 206
Leu Thr Glu Cys Pro Tyr Met Trp Pro Tyr Cys Ser Gln Pro Ile
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Cys Pro Tyr Met Trp Pro Tyr Cys Ser Gln Pro Ile Tyr Tyr Gly
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<211> 1052
<212> PRT
<213> Homo sapiens
<400> 208
Met Lys Leu Val Asn Ile Trp Leu Leu Leu Val Val Leu Leu Cys
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Gly Lys Lys His Leu Gly Asp Arg Leu Glu Lys Lys Ser Phe Glu Lys
Ala Pro Cys Pro Gly Cys Ser His Leu Thr Leu Lys Val Glu Phe Ser
        35 40
Ser Thr Val Val Glu Tyr Glu Tyr Ile Val Ala Phe Asn Gly Tyr Phe
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Glu	Val	Asp	Asn	Trp 85	Arg	Ile	Ile	Pro	Arg 90	Asn	Asn	Pro	Ser	Ser 95	Asp
Tyr	Pro	Ser	Asp 100	Phe	Glu	Val	Ile	Gln 105	Ile	Lys	Glu	Lys	Gln 110	Lys	Ala
Gly	Leu	Leu 115	Thr	Leu	Glu	Asp	His 120	Pro	Asn	Ile	Lys	Arg 125	Val	Thr	Pro
Gln	Arg 130	Lys	Val	Phe	Arg	Ser 135	Leu	Lys	Tyr	Ala	Glu 140	Ser	Asp	Pro	Thr
Val 145	Pro	Cys	Asn	Glu	Thr 150	Arg	Trp	Ser	Gln	Lys 155	Trp	Gln	Ser	Ser	Arg 160
Pro	Leu	Arg	Arg	Ala 165	Ser	Leu	Ser	Leu	Gly 170	Ser	Gly	Phe	Trp	His 175	Ala
Thr	Gly	Arg	His 180	Ser	Ser	Arg	Arg	Leu 185	Leu	Arg	Ala	Ile	Pro 190	Arg	Gln
Val	Ala	Gln 195	Thr	Leu	Gln	Ala	Asp 200	Val	Leu	Trp	Gln	Met 205	Gly	Tyr	Thr
Gly	Ala 210	Asn	Val	Arg	Val	Ala 215	Val	Phe	Asp	Thr	Gly 220	Leu	Ser	Glu	Lys
His 225	Pro	His	Phe	Lys	Asn 230	Val	Lys	Glu	Arg	Thr 235	Asn	Trp	Thr	Asn	Glu 240
Arg	Thr	Leu	Asp	Asp 245	Gly	Leu	Gly	His	Gly 250	Thr	Phe	Val	Ala	Gly 255	Val
Ile	Ala	Ser	Met 260	Arg	Glu	Cys	Gln	Gly 265	Phe	Ala	Pro	Asp	Ala 270	Glu	Leu
His	Ile	Phe 275	Arg	Val	Phe	Thr	Asn 280	Asn	Gln	Val	Ser	Tyr 285	Thr	Ser	Trp
Phe	Leu 290	Asp	Ala	Phe	Asn	Tyr 295	Ala	Ile	Leu	Lys	Lys 300	Ile	Asp	Val	Leu
Asn 305	Leu	Ser	Ile	Gly	Gly 310	Pro	Asp	Phe	Met	Asp 315	His	Pro	Phe	Val	Asp 320
Lys	Val	Trp	Glu	Leu 325	Thr	Ala	Asn	Asn	Val 330	Ile	Met	Val	Ser	Ala 335	Ile
Gly	Asn	Asp	Gly 340	Pro	Leu	Tyr	Gly	Thr 345	Leu	Asn	Asn	Pro	Ala 350	Asp	Gln
Met	Asp	Val	Ile	Gly	Val	Gly	Gly	Ile	Asp	Phe	Glu	Asp	Asn	Ile	Ala

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Arg	Phe 370	Ser	Ser	Arg	Gly	Met 375	Thr	Thr	Trp	Glu	Leu 380	Pro	Gly	Gly	Tyr
Gly 385	Arg	Met	Lys	Pro	Asp 390	Ile	Val	Thr	Tyr	Gly 395	Ala	Gly	Val	Arg	Gly 400
Ser	Gly	Val	Lys	Gly 405	Gly	Cys	Arg	Ala	Leu 410	Ser	Gly	Thr	Ser	Val 415	Ala
Ser	Pro	Val	Val 420	Ala	Gly	Ala	Val	Thr 425	Leu	Leu	Val	Ser	Thr 430	Val	Gln
Lys	Arg	Glu 435	Leu	Val	Asn	Pro	Ala 440	Ser	Met	Lys	Gln	Ala 445	Leu	Ile	Ala
Ser	Ala 450	Arg	Arg	Leu	Pro	Gly 455	Val	Asn	Met	Phe	Glu 460	Gln	Gly	His	Gly
Lys 465	Leu	Asp	Leu	Leu	Arg 470	Ala	Tyr	Gln	Ile	Leu 475	Asn	Ser	Tyr	Lys	Pro 480
Gln	Ala	Ser	Leu	Ser 485	Pro	Ser	Tyr	Ile	Asp 490	Leu	Thr	Glu	Cys	Pro 495	Tyr
Met	Trp	Pro	Туr 500	Cys	Ser	Gln	Pro	Ile 505	Tyr	Tyr	Gly	Gly	Met 510	Pro	Thr
Val	Val	Asn 515	Val	Thr	Ile	Leu	Asn 520	Gly	Met	Gly	Val	Thr 525	Gly	Arg	Ile
Val	Asp 530	Lys	Pro	Asp	Trp	Gln 535	Pro	Tyr	Leu	Pro	Gln 540	Asn	Gly	Asp	Asn
Ile 545	Glu	Val	Ala	Phe	Ser 550	Tyr	Ser	Ser	Val	Leu 555	Trp	Pro	Trp	Ser	Gly 560
Tyr	Leu	Ala	Ile	Ser 565	Ile	Ser	Val	Thr	Lys 570	Lys	Ala	Ala	Ser	Trp 575	Glu
Gly	Ile	Ala	Gln 580	Gly	His	Val	Met	Ile 585	Thr	Val	Ala	Ser	Pro 590	Ala	Glu
Thr	Glu	Ser 595	Lys	Asn	Gly	Ala	Glu 600	Gln	Thr	Ser	Thr	Val 605	Lys	Leu	Pro
Ile	Lys 610	Val	Lys	Ile	Ile	Pro 615	Thr	Pro	Pro	Arg	Ser 620	Lys	Arg	Val	Leu
Trp 625	Asp	Gln	Tyr	His	Asn 630	Leu	Arg	Tyr	Pro	Pro 635	Gly	Tyr	Phe	Pro	Arg 640
Asp	Asn	Leu	Arg	Met 645	Lys	Asn	Asp	Pro	Leu 650	Asp	Trp	Asn	Gly	Asp 655	His
Ile	His	Thr	Asn	Phe	Arg	Asp	Met	Tyr	Gln	His	Leu	Arg	Ser	Met	Gly

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Tyr	Phe	Val 675	Glu	Val	Leu	Gly	Ala 680	Pro	Phe	Thr	Суз	Phe 685	Asp	Ala	Ser
Gln	Тут 690	Gly	Thr	Leu	Leu	Met 695	Val	Asp	Ser	Glu	Glu 700	Glu	Tyr	Phe	Pro
Glu 705	Glu	Ile	Ala	Lys	Leu 710	Arg	Arg	Asp	Val	Asp 715	Asn	Gly	Leu	Ser	Leu 720
Val	Ile	Phe	Ser	Asp 725	Trp	Tyr	Asn	Thr	Ser 730	Val	Met	Arg	Lys	Val 735	Lys
Phe	Τγr	Asp	Glu 740	Asn	Thr	Arg	Gln	Trp 745	Trp	Met	Pro	Asp	Thr 750	Gly	Gly
Ala	Asn	Ile 755	Pro	Ala	Leu	Asn	Glu 760	Leu	Leu	Ser	Val	Trp 765	Asn	Met	Gly
Phe	Ser 770	Asp	Gly	Leu	Tyr	Glu 775	Gly	Glu	Phe	Thr	Leu 780	Ala	Asn	His	Asp
Met 785	Tyr	Tyr	Ala	Ser	Gly 790	Cys	Ser	Ile	Ala	Lys 795	Phe	Pro	Glu	Asp	Gly 800
Val	Val	Ile	Thr	Gln 805	Thr	Phe	Lys	Asp	Gln 810	Gly	Leu	Glu	Val	Leu 815	Lys
Gln	Glu	Thr	Ala 820	Val	Val	Glu	Asn	Val 825	Pro	Ile	Leu	Gly	Leu 830	Tyr	Gln
Ile	Pro	Ala 835	Glu	Gly	Gly	Gly	Arg 840	Ile	Val	Leu	Tyr	Gly 845	Asp	Ser	Asn
Cys	Leu 850	Asp	Asp	Ser	His	Arg 855	Gln	Lys	Asp	Cys	Phe 860	Trp	Leu	Leu	Asp
Ala 865	Leu	Leu	Gln	Tyr	Thr 870	Ser	Tyr	Gly	Val	Thr 875	Pro	Pro	Ser	Leu	Ser 880
His	Ser	Gly	Asn	Arg 885	Gln	Arg	Pro	Pro	Ser 890	Gly	Ala	Gly	Ser	Val 895	Thr
Pro	Glu	Arg	Met 900	Glu	Gly	Asn	His	Leu 905	His	Arg	Tyr	Ser	Lys 910	Val	Leu
Glu	Ala	His 915	Leu	Gly	Asp	Pro	Lys 920	Pro	Arg	Pro	Leu	Pro 925	Ala	Cys	Pro
Arg	Leu 930	Ser	Trp	Ala	Lys	Pro 935	Gln	Pro	Leu	Asn	Glu 940	Thr	Ala	Pro	Ser
Asn 945	Leu	Trp	Lys	His	Gln 950	Lys	Leu	Leu	Ser	Ile 955	Asp	Leu	Asp	Lys	Val 960
Val	Leu	Pro	Asn	Phe	Arg	Ser	Asn	Arg	Pro	Gln	Val	Arg	Pro	Leu	Ser

965 970 975

Pro Gly Glu Ser Gly Ala Trp Asp Ile Pro Gly Gly Ile Met Pro Gly 980 985 990

Arg Tyr Asn Gln Glu Val Gly Gln Thr Ile Pro Val Phe Ala Phe Leu 995 1000 1005

Gly Ala Met Val Val Leu Ala Phe Phe Val Val Gln Ile Asn Lys Ala 1010 1015 1020

Lys Ser Arg Pro Lys Arg Arg Lys Pro Arg Val Lys Arg Pro Gln Leu 1025 1030 1035 1040

Met Gln Gln Val His Pro Pro Lys Thr Pro Ser Val 1045 1050

<210> 209

<211> 280

<212> PRT

<213> Homo sapiens

<400> 209

Arg Ala Ile Pro Arg Gln Val Ala Gln Thr Leu Gln Ala Asp Val Leu 1 5 10 15

Trp Gln Met Gly Tyr Thr Gly Ala Asn Val Arg Val Ala Val Phe Asp 20 25 30

Thr Gly Leu Ser Glu Lys His Pro His Phe Lys Asn Val Lys Glu Arg

Thr Asn Trp Thr Asn Glu Arg Thr Leu Asp Asp Gly Leu Gly His Gly 50 55 60

Thr Phe Val Ala Gly Val Ile Ala Ser Met Arg Glu Cys Gln Gly Phe 65 70 75 80

Ala Pro Asp Ala Glu Leu His Ile Phe Arg Val Phe Thr Asn Asn Gln 85 90 95

Val Ser Tyr Thr Ser Trp Phe Leu Asp Ala Phe Asn Tyr Ala Ile Leu 100 105 110

Lys Lys Ile Asp Val Leu Asn Leu Ser Ile Gly Gly Pro Asp Phe Met 115 120 125

Asp His Pro Phe Val Asp Lys Val Trp Glu Leu Thr Ala Asn Asn Val 130 135 140

Asn Asn Pro Ala Asp Gln Met Asp Val Ile Gly Val Gly Gly Ile Asp 165 170 175

Phe Glu Asp Asn Ile Ala Arg Phe Ser Ser Arg Gly Met Thr Thr Trp 180 185 Glu Leu Pro Gly Gly Tyr Gly Arg Met Lys Pro Asp Ile Val Thr Tyr 200 Gly Ala Gly Val Arg Gly Ser Gly Val Lys Gly Gly Cys Arg Ala Leu 210 215 Ser Gly Thr Ser Val Ala Ser Pro Val Val Ala Gly Ala Val Thr Leu 230 235 Leu Val Ser Thr Val Gln Lys Arg Glu Leu Val Asn Pro Ala Ser Met 250 Lys Gln Ala Leu Ile Ala Ser Ala Arg Arg Leu Pro Gly Val Asn Met 270 260 265 Phe Glu Gln Gly His Gly Lys Leu 275 <210> 210 <211> 15 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic <400> 210 Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val 5 10 <210> 211 <211> 15 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic <400> 211 Ala Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val <210> 212 <211> 15 <212> PRT <213> Artificial Sequence <223> Description of Artificial Sequence: Synthetic

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Gly Ser Ile Ala Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val
<210> 215
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<213> Artificial Sequence
<223> Description of Artificial Sequence: Synthetic
<400> 215
Gly Ser Ile Ser Ala Pro Ala Arg Tyr Ala Asn Ala Met Ala Val
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<213> Artificial Sequence
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Gly Ser Ile Ser Tyr Ala Ala Arg Tyr Ala Asn Ala Met Ala Val
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Gly Ser Ile Ser Tyr Pro Ala Ala Tyr Ala Asn Ala Met Ala Val
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<400> 218
Gly Ser Ile Ser Tyr Pro Ala Arg Ala Ala Asn Ala Met Ala Val
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Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Ala Ala Met Ala Val
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Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Ala Ala Val
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<211> 15

<212> PRT

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<400> 222

Pro Gly Gly Val Ala Tyr Ser Cys Ala Asp Gln Thr Pro Trp Ala 1 5 10 15

<210> 223

<211> 15

<212> PRT

<213> Humicola insolens

<400> 223

Cys Gly Trp Ala Lys Lys Ala Pro Val Asn Gln Pro Val Phe Ser
1 5 10 15

<210> 224

<211> 276

<212> PRT

<213> Humicola insolens

<400> 224

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Val Leu Ala Leu Ala Asp Gly Arg Ser Thr Arg Tyr Trp Asp Cys
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Cys Lys Pro Ser Cys Gly Trp Ala Lys Lys Ala Pro Val Asn Gln Pro 35 40 45

Val Phe Ser Cys Asn Ala Asn Phe Gln Arg Ile Thr Asp Phe Asp Ala 50 55 60

Lys Ser Gly Cys Glu Pro Gly Gly Val Ala Tyr Ser Cys Ala Asp Gln 65 70 75 80

Thr Pro Trp Ala Val Asn Asp Asp Phe Ala Leu Gly Phe Ala Ala Thr 85 90 95

Ser Ile Ala Gly Ser Asn Glu Ala Gly Trp Cys Cys Ala Cys Tyr Glu 100 105 110

Leu Thr Phe Thr Ser Gly Pro Val Ala Gly Lys Lys Met Val Val Gln
115 120 125

Ser Thr Ser Thr Gly Gly Asp Leu Gly Ser Asn His Phe Asp Leu Asn 130 135 140

Ile Pro Gly Gly Gly Val Gly Ile Phe Asp Gly Cys Thr Pro Gln Phe 145 150 155 160

Gly Gly Leu Pro Gly Gln Arg Tyr Gly Gly Ile Ser Ser Arg Asn Glu 165 170 175

Cys Asp Arg Phe Pro Asp Ala Leu Lys Pro Gly Cys Tyr Trp Arg Phe 180 185 190

Asp Trp Phe Lys Asn Ala Asp Asn Pro Ser Phe Ser Phe Arg Gln Val 195 200 205

Gln Cys Pro Ala Glu Leu Val Ala Arg Thr Gly Cys Arg Arg Asn Asp 210 215 220

Asp Gly Asn Phe Pro Ala Val Gln Ile Pro Ser Ser Ser Thr Ser Ser 225 230 235 240

Pro Val Asn Gln Pro Thr Ser Thr Ser Thr Thr Ser Thr Ser Thr Thr 245 250 255

Ser Ser Pro Pro Val Gln Pro Thr Thr Pro Ser Gly Cys Thr Ala Glu 260 265 270

Arg Trp Ala Gln 275

<210> 225

<211> 18

<212> PRT

<213> Thermomyces lanuginosus

<400> 225

Gly Asp Val Thr Gly Phe Leu Ala Leu Asp Asn Thr Asn Lys Leu Ile 1 5 10 15

Val Leu

<210> 226

<211> 15

<212> PRT

<213> Thermomyces lanuginosus

<400> 226

Ser Ile Glu Asn Trp Ile Gly Asn Leu Asn Phe Asp Leu Lys Glu
1 10 15

<210> 227

<211> 291

<212> PRT

<213> Thermomyces lanuginosus

<400> 227

- Met Arg Ser Ser Leu Val Leu Phe Phe Val Ser Ala Trp Thr Ala Leu 1 5 10 15
- Ala Ser Pro Ile Arg Arg Glu Val Ser Gln Asp Leu Phe Asn Gln Phe 20 25 30
- Asn Leu Phe Ala Gln Tyr Ser Ala Ala Ala Tyr Cys Gly Lys Asn Asn 35 40 45
- Asp Ala Pro Ala Gly Thr Asn Ile Thr Cys Thr Gly Asn Ala Cys Pro 50 55 60
- Glu Val Glu Lys Ala Asp Ala Thr Phe Leu Tyr Ser Phe Glu Asp Ser
  65 70 75 80
- Gly Val Gly Asp Val Thr Gly Phe Leu Ala Leu Asp Asn Thr Asn Lys
  85 90 95
- Leu Ile Val Leu Ser Phe Arg Gly Ser Arg Ser Ile Glu Asn Trp Ile 100 105 110
- Gly Asn Leu Asn Phe Asp Leu Lys Glu Ile Asn Asp Ile Cys Ser Gly
  115 120 125
- Cys Arg Gly His Asp Gly Phe Thr Ser Ser Trp Arg Ser Val Ala Asp 130 135 140
- Thr Leu Arg Gln Lys Val Glu Asp Ala Val Arg Glu His Pro Asp Tyr 145 150 155 160
- Arg Val Val Phe Thr Gly His Ser Leu Gly Gly Ala Leu Ala Thr Val
- Ala Gly Ala Asp Leu Arg Gly Asn Gly Tyr Asp Ile Asp Val Phe Ser 180 185 190
- Tyr Gly Ala Pro Arg Val Gly Asn Arg Ala Phe Ala Glu Phe Leu Thr
  195 200 205
- Val Gln Thr Gly Gly Thr Leu Tyr Arg Ile Thr His Thr Asn Asp Ile 210 215 220
- Val Pro Arg Leu Pro Pro Arg Glu Phe Gly Tyr Ser His Ser Ser Pro 225 230 235 240
- Glu Tyr Trp Ile Lys Ser Gly Thr Leu Val Pro Val Thr Arg Asn Asp 245 250 255
- Ile Val Lys Ile Glu Gly Ile Asp Ala Thr Gly Gly Asn Asn Gln Pro 260 265 270
- Asn Ile Pro Asp Ile Pro Ala His Leu Trp Tyr Phe Gly Leu Ile Gly 275 280 285

Thr Cys Leu 290

<210> 228

<211> 15

<212> PRT

<213> Streptomyces plicatus

<400> 228

Ile Lys Val Leu Leu Ser Val Leu Gly Asn His Gln Gly Ala Gly
1 5 10 15

<210> 229

<211> 313

<212> PRT

<213> Streptomyces plicatus

<400> 229

Met Phe Thr Pro Val Arg Arg Arg Val Arg Thr Ala Ala Leu Ala Leu 1 5 10 15

Ser Ala Ala Ala Leu Val Leu Gly Ser Thr Ala Ala Ser Gly Ala
20 25 30

Ser Ala Thr Pro Ser Pro Ala Pro Ala Pro Ala Pro Ala Pro Val Lys
35 40 45

Gln Gly Pro Thr Ser Val Ala Tyr Val Glu Val Asn Asn Asn Ser Met 50 55 60

Leu Asn Val Gly Lys Tyr Thr Leu Ala Asp Gly Gly Asn Ala Phe 65 70 75 80

Asp Val Ala Val Ile Phe Ala Ala Asn Ile Asn Tyr Asp Thr Gly Thr 85 90 95

Lys Thr Ala Tyr Leu His Phe Asn Glu Asn Val Gln Arg Val Leu Asp 100 105 110

Asn Ala Val Thr Gln Ile Arg Pro Leu Gln Gln Gln Gly Ile Lys Val 115 120 125

Leu Leu Ser Val Leu Gly Asn His Gln Gly Ala Gly Phe Ala Asn Phe 130 135 140

Pro Ser Gln Gln Ala Ala Ser Ala Phe Ala Lys Gln Leu Ser Asp Ala 145 150 155 160

Val Ala Lys Tyr Gly Leu Asp Gly Val Asp Phe Asp Asp Glu Tyr Ala 165 170 175

Glu Tyr Gly Asn Asn Gly Thr Ala Gln Pro Asn Asp Ser Ser Phe Val

His Leu Val Thr Ala Leu Arg Ala Asn Met Pro Asp Lys Ile Ile Ser

195 200 205

Leu Tyr Asn Ile Gly Pro Ala Ala Ser Arg Leu Ser Tyr Gly Gly Val 210 215 220

Asp Val Ser Asp Lys Phe Asp Tyr Ala Trp Asn Pro Tyr Tyr Gly Thr 225 230 235 240

Trp Gln Val Pro Gly Ile Ala Leu Pro Lys Ala Gln Leu Ser Pro Ala 245 250 255

Ala Val Glu Ile Gly Arg Thr Ser Arg Ser Thr Val Ala Asp Leu Ala 260 265 270

Arg Arg Thr Val Asp Glu Gly Tyr Gly Val Tyr Leu Thr Tyr Asn Leu 275 280 285

Asp Gly Gly Asp Arg Thr Ala Asp Val Ser Ala Phe Thr Arg Glu Leu 290 295 300

Tyr Gly Ser Glu Ala Val Arg Thr Pro 305 310

<210> 230

<211> 15

<212> PRT

<213> Bacillus amyloliquefaciens

<400> 230

Gly Thr Val Ala Ala Leu Asn Asn Ser Ile Gly Val Leu Gly Val
1 5 10 15

<210> 231

<211> 15

<212> PRT

<213> Bacillus amyloliquefaciens

<400> 231

Asn Gly Ile Glu Trp Ala Ile Ala Asn Asn Met Asp Val Ile Asn 1 5 10 15

<210> 232

<211> 15

<212> PRT

<213> Bacillus lentus

<400> 232

Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp Thr Gly Ile Ser
1 5 10 15

<210> 233

<211> 15

<212> PRT

<213> Bacillus lentus

<400> 233

Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala Ser Gly Ser 1 5 10 15

<210> 234

<211> 17

<212> PRT

<213> Bacillus lentus

<400> 234

Gly Ser Ile Ser Tyr Pro Ala Arg Tyr Ala Asn Ala Met Ala Val Gly
1 5 10 15

Ala

<210> 235

<211> 15

<212> PRT

<213> Bacillus lentus

<400> 235

Gly Ala Gly Leu Asp Ile Val Ala Pro Gly Val Asn Val Gln Ser
1 5 10 15

<210> 236

<211> 272

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Hybrid of Bacillus lentus and Bacillus amyloliquefaciens

<400> 236

Ala Gln Ser Val Pro Trp Gly Ile Ser Arg Val Gln Ala Pro Ala Ala 1 5 10 15

His Asn Arg Gly Leu Thr Gly Ser Gly Val Lys Val Ala Val Leu Asp 20 25 30

Thr Gly Ile Ser Thr His Pro Asp Leu Asn Ile Arg Gly Gly Ala Ser 35 40 45

Phe Val Pro Gly Glu Pro Ser Thr Gln Asp Gly Asn Gly His Gly Thr 50 55 60

His Val Ala Gly Thr Ile Ala Ala Leu Asn Asn Ser Ile Gly Val Leu 65 70 75 80

Gly Val Ala Pro Ser Ala Glu Leu Tyr Ala Val Lys Val Leu Gly Ala 85 90 95

Ser Gly Ser Gly Ser Val Ser Ser Ile Ala Gln Gly Leu Glu Trp Ala 105 Gly Asn Asn Gly Met His Val Ile Asn Met Ser Leu Gly Gly Ser Gly Ser Ala Ala Leu Lys Ala Ala Val Asp Lys Ala Val Ala Ser Gly Val 135 Val Val Val Ala Ala Ala Gly Asn Glu Gly Thr Ser Gly Ser Ser 150 155 Thr Val Gly Tyr Pro Gly Lys Tyr Pro Ser Val Ile Ala Val Gly Ala 165 170 Val Asp Ser Ser Asn Gln Arg Ala Ser Phe Ser Ser Val Gly Pro Glu 185 Leu Asp Val Met Ala Pro Gly Val Ser Ile Gln Ser Thr Leu Pro Gly 195 Asn Lys Tyr Gly Ala Tyr Asn Gly Thr Ser Met Ala Ser Pro His Val 215 Ala Gly Ala Ala Leu Ile Leu Ser Lys His Pro Asn Trp Thr Asn 230 235 Thr Gln Val Arg Ser Ser Leu Glu Asn Thr Thr Thr Lys Leu Gly Asp 245 250 Ser Phe Tyr Tyr Gly Lys Gly Leu Ile Asn Val Gln Ala Ala Gln 265